

IN THE SPECIFICATION

Please delete the entire Table 1, as well as the five lines of commentary underneath this Table at pages 78 and 79 of the specification and replace it with the following Table 1 and commentary:

TABLE 1

Summary of the protein-protein interactions
between the SID polypeptides of the invention
and H77 strain HCV polypeptides

Bait	SED ID NO (1)	begin (2)	end (2)	SEQ ID NO (3)	SID	SEQ ID NO (4)	begin (2)	End (2)	SEQ ID NO (5)
Core (87%)	114	302	614	77	Core (100%)	39	446	600	1
Core (100%)	115	342	683	78	NS3 (100%)	40	4814	4922	2
Core (100%)	115	342	683	78	Core (100%)	41	380	616	3
E1 (100%)	116	995	1342	79	E2 (100%)	42	1871	1987	4
E1 (4%) / E2 (95%)	117	1478	1756	80	NS3 (100%)	43	4787	5242	5
E2 (100%)	118	1745	2278	81	E2 (100%)	44	1871	1958	6
E2 (100%)	119	1799	2090	82	E2 (100%)	45	1808	1890	7
NS2 (12%) / NS3 (87%)	120	3312	4150	83	NS4A (59%) / NS4B (40%)	46	5375	5542	8
NS3 (100%)	121	3767	4244	84	NS3 (100%)	47	4676	4801	9
NS3 (100%)	122	3779	4571	85	NS3 (100%)	48	4856	4945	10
NS3 (100%)	123	3974	4559	86	NS3 (100%)	49	4817	4903	11
NS3 (100%)	124	4238	4857	87	NS5B (100%)	50	7979	8109	12
NS3 (100%)	125	4298	4859	88	NS3 (100%)	51	4031	4118	13
NS3 (100%)	126	4691	5168	89	E2 (100%)	52	1784	1888	14
NS3 (100%)	127	4838	5230	90	E2 (100%)	53	1871	1968	15
NS3 (1%) / NS4A (98%)	128	5310	5467	91	NS4B (100%)	54	5918	6154	16
NS4A (100%)	129	5342	5400	92	NS3 (100%)	55	3512	3956	17
NS4B (86%) /NS5A (13%)	130	5717	6344	93	NS4B (53%) / NS5A (46%)	56	6197	6310	18
NS4B (70%) /NS5A (29%)	131	5819	6444	94	E2 (100%)	57	1844	1933	19

NS4B (55%) /NS5A (44%)	132	5882	6562	95	NS5B (100%)	58	9083	9222	20
NS4B (82%) / NS5A (17%)	133	5897	6335	96	NS4B (100%)	59	5819	6080	21
NS4B (100%)	134	6011	6177	97	E2 (100%)	60	1823	1955	22

TABLE 1 (continued)

Bait	SED ID NO (1)	begin (2)	end (2)	SEQ ID NO (3)	SID	SEQ ID NO (4)	begin (2)	End (2)	SEQ ID NO (5)
NS4B (30%) / NS5A (69%)	135	6107	6605	98	NS4B (100%)	61	5879	6072	23
NS4B (12%) / NS5A (87%)	136	6141	7069	99	E2 (100%)	62	1784	1875	24
NS4B (8%) / NS5A (91%)	137	6182	7034	100	E1 (100%)	63	1226	1458	25
NS4B (9%) / NS5A (90%)	138	6188	6939	101	NS4B (70%) / NS5A (28%)	64	6176	6291	26
NS5A (100%)	139	6317	6576	102	NS3 (100%)	65	4784	4928	27
NS5A (100%)	140	6440	6727	103	NS5A (100%)	66	6557	6721	28
NS5A (100%)	141	7019	7249	104	NS3 (100%)	67	4451	4790	29
NS5A (100%)	142	7274	7549	105	NS4B (100%)	68	6029	6194	30
NS5B (100%)	143	7613	8027	106	NS5B (100%)	69	8354	8665	31
NS5B (100%)	144	7838	8743	107	NS5B (100%)	70	7769	8011	32
NS5B (100%)	145	7856	8458	108	NS3 (100%)	71	4715	4901	33
NS5B (100%)	146	7976	8759	109	NS5B (100%)	72	7775	8011	34
NS5B (100%)	147	8564	8948	110	E2 (100%)	73	1805	1887	35
NS5B (100%)	148	8708	8978	111	E2 (100%)	74	1751	1865	36
NS5B (100%)	149	8996	9220	112	NS4B (57%) / NS5A (41%)	75	6194	6303	37
NS5B (100%)	150	9032	9226	113	NS4B (63%) / NS5A (35%)	76	6206	6286	38

(1) Nucleic acid sequence encoding the polypeptide from the H77 strain of HCV, which binds to the SID polypeptide (4) described in the same line.

(2) 5'-end and 3'-end nucleotide positions of the sequence SED ID (1) in reference to the nomenclature disclosed by Yanagi et al. (1997)

(3) Amino acid sequence of the polypeptide from the H77 strain of HCV, which binds to the SID polypeptide (4) described in the same line.

(4) Nucleic acid sequence encoding the SID polypeptide, which binds to the polypeptide of the amino acid sequence (3) described in the same line.

(5) Amino acid sequence of the SID polypeptide, which binds to the polypeptide of the amino acid sequence (3) described in the same line.